

#8

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PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: PCT/US02/16792

DATE: 06/11/2002
TIME: 13:14:02

Input Set : A:\555258057W00-seqlist.txt
Output Set: N:\CRF3\06112002\PU16792.raw

ENTERED

4 <110> APPLICANT: Lynx Therapeutics, Inc.
6 <120> TITLE OF INVENTION: Sequencing by Proxy
8 <130> FILE REFERENCE: 555258057W00
10 <140> CURRENT APPLICATION NUMBER: PCT/US02/16792
11 <141> CURRENT FILING DATE: 2002-05-29
13 <150> PRIOR APPLICATION NUMBER: US 09/867,201
14 <151> PRIOR FILING DATE: 2001-05-29
16 <160> NUMBER OF SEQ ID NOS: 4
18 <170> SOFTWARE: FastSEQ for Windows Version 4.0
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 65
22 <212> TYPE: DNA
23 <213> ORGANISM: Artificial Sequence
25 <220> FEATURE:
26 <223> OTHER INFORMATION: exemplary tag library
28 <221> NAME/KEY: misc_feature
29 <222> LOCATION: (1)...(65)
30 <223> OTHER INFORMATION: n = A,T,C or G
32 <400> 1
33 agaattcggg ccttaattaa dddddddddd dddggggccc gcataagtct 60
34 tcnnn 65
36 <210> SEQ ID NO: 2
37 <211> LENGTH: 16
38 <212> TYPE: DNA
39 <213> ORGANISM: Artificial Sequence
41 <220> FEATURE:
42 <223> OTHER INFORMATION: 3' end of exemplary tag sequence
44 <221> NAME/KEY: misc_feature
45 <222> LOCATION: (1)...(16)
46 <223> OTHER INFORMATION: n = A,T,C or G
48 <400> 2
49 nnnnggatccg agtgat 16
51 <210> SEQ ID NO: 3
52 <211> LENGTH: 20
53 <212> TYPE: DNA
54 <213> ORGANISM: Artificial Sequence
56 <220> FEATURE:
57 <223> OTHER INFORMATION: primer
59 <400> SEQUENCE: 3
60 agaattcggg ccttaattaa 20
62 <210> SEQ ID NO: 4
63 <211> LENGTH: 18
64 <212> TYPE: DNA

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65 <213> ORGANISM: Artificial Sequence

67 <220> FEATURE:

68 <223> OTHER INFORMATION: primer

W--> 70 <221> NAME/KEY: misc_feature

71 <222> LOCATION: (1)...(18)

72 <223> OTHER INFORMATION: n = A,T,C or G

W--> 74 <400> 4

W--> 75 nnnnncctag gctcacta

18

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: PCT/US02/16792

DATE: 06/11/2002
TIME: 13:14:03

Input Set : A:\555258057W00-seqlist.txt
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 63,64,65
Seq#:2; N Pos. 1,2,3
Seq#:4; N Pos. 1,2,3,4,5